

W098/18930 SEQ ID 66

Query Match
Best Local Similarity 80.9%; Score 575.5; DB 4; Length 763;
Matches 103; Conservative 18; Mismatches 9; Indels 1; Gaps 1;
7 CAYALNQHRS-GENKDNRRVSYVDGSSQSKSENLTTPDOVSQKEGIAEQIIVIKITTDGY 65
1 CSYELGRHQAQGVKKESNRVSIIDGDAQAKAENLTPDEVSKREGINAEQXVIRIKITTDGY 60
66 VTSHGDHYYHYNGKVPYDALFSEELLTKKDPNYYQLKADADIVNEVKGGYIIIKVDGKYYVYLK 125
61 VTSHGDHYYHYNGKVPYDALFSEELLTKKDPNYYQLKADSDIVNEIKGKGYVIRKNGKYYVYLK 120
126 DAHADNVRTK 136
121 DAHADNIRTK 131

W098/18930 SEQ ID 56

Query Match
Best Local Similarity 80.7%; Score 574; DB 4; Length 796;
Matches 104; Conservative 16; Mismatches 9; Indels 0; Gaps 0;
8 AYALNQHRSQENKDNRRVSYVDGSSQSKSENLTTPDOVSQKEGIAEQIIVIKITTDGY 67
1 SYELGLYQARTVKENNRVSIIDGKQATOTKTENLTPDEVSKREGINAEQIIVIKITTDGY 60
68 SHGDHYYHYNGKVPYDALFSEELLTKKDPNYYQLKADADIVNEVKGGYIIIKVDGKYYVYLK 127
61 SHGDHYYHYNGKVPYDALFSEELLTKKDPNYYQLKADSDIVNEIKGKGYVIRKNGKYYVYLK 120
128 AHADNVRTK 136
121 AHADNVRTK 129

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7 CAYALNQHRS-GENKDNRRVSYVDGSSSSQKSENLTTPDOVSQKREGIOAEQIVIKITIDGY 65
1 CSYELGRHQAAGQVKKESNRVSYIDGDOAGOKAENLTTPDEVSKREGINAEQIXYIKITIDGY 60
66 VTSHGDDHYHYNGKVPYDALFSEELTMKDPNYYQLKADADIVNEVKGGYIIKVDGKXYVYLK 125
61 VTSHGDDHYHYNGKVPYDALFSEELTMKDPNYYQLKADSDIVNEIKGGYIVIKVNGKXYVYLK 120
DB 61 VTSHGDDHYHYNGKVPYDALFSEELTMKDPNYYQLKADSDIVNEIKGGYIVIKVNGKXYVYLK 120
OY 126 DAHADNVRTK 136
DB 121 DAHADNVRTK 131

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Query Match
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Matches 104; Conservative 16; Mismatches 9; Indels 0; Gaps 0;
8 AYALNQHRS-GENKDNRRVSYVDGSSSSQKSENLTTPDOVSQKREGIOAEQIVIKITIDGY 67
1 SYELGLYQARTVKKENNRVSYIDGKQATQKTENLTTPDEVSKREGINAEQIVIKITIDGY 60
68 SHGDHYHYHYNGKVPYDALFSEELTMKDPNYYQLKADADIVNEVKGGYIIKVDGKXYVYLK 127
61 SHGDHYHYHYNGKVPYDALFSEELTMKDPNYYQLKADSDIVNEVKGGYIVIKVNGKXYVYLK 120
DB 61 SHGDHYHYHYNGKVPYDALFSEELTMKDPNYYQLKADSDIVNEVKGGYIVIKVNGKXYVYLK 120
OY 128 AHADNVRTK 136
DB 121 AHADNVRTK 129

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